

**In the specification:**

Please amend the paragraph beginning on page 18, line 35 and continuing to page 19, lines 1-2, as follows:

***In silico* analysis of T128**

The amino acid sequence of T128 was analysed using search programs including PROSITE (~~accessible at [www.expasy.ch/prosite/](http://www.expasy.ch/prosite/)~~), PSORT (~~accessible at <http://psort.nibb.ac.jp>~~) and Pfam (~~accessible at <http://www.sanger.ac.uk/cgi-bin/Pfam/nph-search.cgi>~~). These identified:

In each instance- on page 2, line 32; page 3, line 4; and page 20, line 29, please replace “Genbank” with -- GenBank® --.

On page 17, line 35, and also on page 18, line 18, please replace “Mx4000” with -- Mx4000® --.